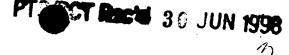
## SEQUENCE LISTING



## (1) GENERAL INFORMATION:

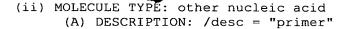
- (i) APPLICANT: Klatzmann, David Cohen, Jacques
- (ii) TITLE OF INVENTION:  $\alpha \beta$  C4BP-TYPE RECOMBINANT HETEROMULTIMERIC PROTEINS
- (iii) NUMBER OF SEQUENCES: 6
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: SPENCER & FRANK
  - (B) STREET: 1100 New York Avenue, N.W., Suite 300 East
  - (C) CITY: Washington
  - (D) STATE: DC
  - (E) COUNTRY: USA
  - (F) ZIP: 20005
  - (v) COMPUTER READABLE FORM:
    - (A) MEDIUM TYPE: Floppy disk
    - (B) COMPUTER: IBM PC compatible
    - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
    - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: PCT/FR96/01132
  - (B) FILING DATE: 18-JUL-1996
  - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: Gollin, Michael A.
  - (B) REGISTRATION NUMBER: 31,957
  - (C) REFERENCE/DOCKET NUMBER: GUPLA 0007
  - (ix) TELECOMMUNICATION INFORMATION:
    - (A) TELEPHONE: 202-414-4000
    - (B) TELEFAX: 202-414-4040
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 20 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: other nucleic acid
    - (A) DESCRIPTION: /desc = "primer"
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GAGACCCCCG AAGGCTGTGA

(2) INFORMATION FOR SEQ ID NO:2:

20

	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 29 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "primer"	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:	
CTC	GAGTT	AT AGTTCTTTAT CCCAAGTGG	29
(2)	INFO	RMATION FOR SEQ ID NO:3:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 28 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "primer"	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GCG	GCCGC	AG AGACCCCGA AGGCTGTG	28
(2)	INFO	RMATION FOR SEQ ID NO:4:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 23 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "primer"	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:4:	
CCA	CTTTG	GA TAAAGAACTA TAA	23
(2)	INFO	RMATION FOR SEQ ID NO:5:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 36 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	



(xi)	SEQUENCE	DESCRIPTION:	SEQ	ΙD	NO:5:

36

## (2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

ATTTCTAGAG AGTTATAGTT CTTTATCCAA AGTGGA

- (A) LENGTH: 46 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
   (A) DESCRIPTION: /desc = "primer"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
  CCGGGACAGG TCCTGCTGGA ATCCAACATC AAGGTTCTGC CCACAG

46